

**Amendments to the Specification**

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On page 2, please replace paragraph 3, with the following rewritten paragraph:

The subject specification contains amino acid sequence information prepared using the ~~programme~~ program PatentIn Version 2.0, presented herein after the bibliography. Each amino acid sequence is identified in the sequence listing by the numeric indicator <210> followed by the sequence identifier (e.g. <210>1, <210>2, etc.). The length, type of sequence (protein (PRT), etc) and source organism for each amino acid sequence are indicated by information provided in the numeric indicator fields <211>, <212> and <213>, respectively. ~~Amino acid sequences referred to in the specification are defined by the information provided in numeric indicator field <400> followed by the sequence identifier (e.g. <400>1, <400>2, etc).~~ Amino acid sequences referred to in the specification are identified by the indicator SEQ ID NO: followed by the sequence identifier (eg SEQ ID NO: 1, SEQ ID NO:2, etc). The sequence identifier referred to in the specification correlates to the information provided in numeric indicator field <400> in the sequence listing, which is followed by the sequence identifier (e.g. <400>1, <400>2, etc). That is, SEQ ID NO:1 as detailed in the specification correlates to the sequence indicated as <400>1 in the sequence listing.

Attached is a letter relating to the substitute specification that was provided in Paper no. 28. That substitute specification corresponded to the marked-up version submitted with the Sequence Listing submitted October 15, 2002.